STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/525,/80
Source:	1 FWP
Date Processed by STIC:	1/16/07

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
 Alexandria, VA 22314

Revised 01/10/06

ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/525, /80
ATTN: NEW RULES CASES:	PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3Misaligned Amino / Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)
11Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules
PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid





IFWP

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/525,180

DATE: 01/16/2007

TIME: 15:08:55

Input Set : N:\SSLM\10525180.txt

Output Set: N:\CRF4\01162007\J525180.raw

Use English in a U.S. application

ESEQUENZPROTOROLL SEQUENCE LISTING 3 <110> APPLICANT: Charite Universitatsmedizin Berlin

5 <120> TITLE OF INVENTION: Immunmarker zur Diagnostik and Therapie im Zusammenhang mit

Tranplantat-Reaktionen

8 <130> FILE REFERENCE: P153902PC-La

C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/525,180

C--> 11 <141> CURRENT FILING DATE: 2005-02-22

13 <150> PRIOR APPLICATION NUMBER: DE 102 38 922.5

14 <151> PRIOR FILING DATE: 2002-08-22

16 <160> NUMBER OF SEQ ID NOS: 8

18 <170> SOFTWARE: Patentln Ver. 2.1

see tem 4 on Euro

Does Not Comply Corrected Dickette Needed

ERRORED SEQUENCES

65 <210> SEQ ID NO: 4

66 <211> LENGTH: 310

67 <212> TYPE: DNA

68 <213> ORGANISM: Rattus rattus

70 <400> SEQUENCE: 4

71 acattcatta ttaaatgtga taatagaggt agaggtataa ataatatgaa ggggtgaggg 60

72 aaccagttct acccggtttg ttttgaatgc ttaaattatg taatttaaat agataatctt 120

E--> 73 tacttatgta ggtcttttgg aaata actt ataaatttaa cacagaggac tactactaaa 180

/> 74 cgtgagaggt atgataatcg gcatggaagt tgggctggtt gaccaccaaa gttcaattct 240

75 taaagacatc ttaatcctga atataaaaat gcctttgtgg gtttagaatt agaatttaat 300

76 tttggcattt 310

122 <210> SEO ID NO: 8

123 <211> LENGTH: 313

124 <212> TYPE: DNA

125 <213> ORGANISM: Rattus rattus

127 <400> SEQUENCE: 8

128 aggctagggc tagttctgcg gaccctctcg gagagaggaa taaggttgaa ctgcctgtcc 60

129 ggttctcctt cccctattcc cagatgcagg tggaagcctc cctctagtcc ttccccctaa 120

130 ccgcgacgaa gaccttggct aacacttgct cctttcgcac accatagaaa atgcagtgca 180

131 gacaaacaca gcctcgtcag gcgcttgagg agcgaagtcc aatctgggtc ggcacctgca 240

132 ccaggtettt gegeacetgg teagaagace ggeacecaat agttgettat taaactetae 300

133 gtttgtcccg aaa

313

E--> 134

nuclui nuclui ocid designator

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/525,180

DATE: 01/16/2007 TIME: 15:08:56

Input Set : N:\SSLM\10525180.txt

Output Set: N:\CRF4\01162007\J525180.raw

L:1 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION:

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:73 M:254 E: No. of Bases conflict, LENGTH:Input:180 Counted:181 SEQ:4

L:73 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1

M:254 Repeated in SeqNo=4

L:76 M:252 E: No. of Seq. differs, <211> LENGTH:Input:310 Found:311 SEQ:4

L:134 M:254 E: No. of Bases conflict, this line has no nucleotides.

SEQUENZPROTOKOLL

<110> Charite Universitatsmedizin Berlin

<120> Immunmarker zur Diagnostik and Therapie im Zusammenhang mit Tranplantat-Reaktionen

<130> P153902PC-La

<1507 (1407) PCT/EP03/09355 <1517 (1415) 2003-08-22

Klese are prior application data